Refine Search

Search Results -

Term	Documents
MYCOPLASMA	11249
MYCOPLASMAS	1365
(8 AND MYCOPLASMA).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	15
(L8 AND MYCOPLASMA).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	15

Database: US Pre-Grant Publication Full-Text Database; US Patents Full-Text Database; US OCR Full-Text Database; EPO Abstracts Database; JPO Abstracts Database; Derwent World Patents Index; IBM Technical Disclosure Bulletins

Search: L23

Refine Search | Recall Text | Clear | Interrupt

Search History

DATE: Thursday, March 22, 2007 **Purge Queries** Printable Copy Create Case

Set Name side by side	Query	Hit Count	Set Name result set
DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=OR			
L23	l8 and mycoplasma	15	L23
L22	L21 and mycoplasma	1	L22
L21	l5 or l6	389	L21
L20	l4 and ribosomal adj ribonucleic adj acid	0	L20
L19	l18 and ribosomal RNA	167141	L19
L18	L17 and template	2893	L18
L17	L16 and mixture	3515	L17
L16	L15 and detect$	3821	L16
L15	L14 and hybridi$	3866	L15
L14	L13 and primer	4603	L14

L13	L12 and mycoplasma	9686	L13
L12	L4 and polymerase chain reaction	2952816	L12
L11	l4 and mycoplasma	3	L11
L10	l4 and l6 and l7 and l8	0	L10
L9	l4 and l5 and l6 and l7 and l8	0	L9
L8	dubois.in.	3598	L8
L7	bozyan.in.	3	L7
L6	brown-j.in.	323	L6
L5	brown-j-t.in.	66	L5
L4	happe.in.	294	L4
DB=USPT; PLUR=YES; OP=OR			
L3	6977164.pn.	1	L3
L2	6395470.pn.	1	L2
DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=OR			
L1	6958390.pn.	2	L1

END OF SEARCH HISTORY

STIC REPORT

GenCore version 6.2

OM nucleic - nucleic search, using sw model

Run on: March 22, 2007, 12:31:08 ; Search time 2668 Seconds
(without alignments)
544.055 Million cell updates/sec

Title: US-10-820-971-3
Perfect score: 21
Sequence: 1 aataagccccggctaactatg 21

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 7568541 seqs, 34560148153 residues

Total number of hits satisfying chosen parameters: 15137082

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:*
1: gb_env:*
2: gb_pat:*
3: gb_ph:*
4: gb_pl:*
5: gb_pr:*
6: gb_ro:*
7: gb_sts:*
8: gb_sy:*
9: gb_un:*
10: gb_vi:*
11: gb_ov:*
12: gb_htg:*
13: gb_in:*
14: gb_om:*

```
            Kingsbury,D.T.
  TITLE     Phylogeny of mycoplasmalike organisms (phytoplasmas):
a basis for
            their classification
  JOURNAL   J. Bacteriol. 176 (17), 5244-5254 (1994)
   PUBMED   8071198
COMMENT     Original source text: Blueberry stunt mycoplasma-like
organism DNA.
FEATURES             Location/Qualifiers
     source          1. .339
                     /organism="Blueberry stunt phytoplasma"
                     /mol_type="genomic DNA"
                     /db_xref="taxon:37695"
                     /lab_host="Catharanthus roseus"
ORIGIN

  Query Match             100.0%;  Score 21;  DB 15;  Length 339;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches   21;  Conservative    0;  Mismatches    0;  Indels
0;  Gaps    0;

Qy          1 AATAAGCCCCGGCTAACTATG 21
              |||||||||||||||||||||
Db        314 AATAAGCCCCGGCTAACTATG 334


RESULT 4
MPLRGDPA01
LOCUS       MPLRGDPA01               339 bp    DNA     linear
BCT 21-APR-1995
DEFINITION  Paulownia witches'-broom mycoplasmalike organism 16S
ribosomal RNA
            (16S rRNA) gene, partial sequence.
ACCESSION   L27032
VERSION     L27032.1  GI:780101
KEYWORDS    16S ribosomal RNA.
SEGMENT     1 of 2
SOURCE      Paulownia witches'-broom phytoplasma
  ORGANISM  Paulownia witches'-broom phytoplasma
            Bacteria; Firmicutes; Mollicutes; Acholeplasmatales;
            Acholeplasmataceae; Candidatus Phytoplasma; Candidatus
Phytoplasma
            asteris.
REFERENCE   1  (bases 1 to 339)
  AUTHORS   Gundersen,D.E., Lee,I.M., Rehner,S.A., Davis,R.E. and
            Kingsbury,D.T.
```

```
  TITLE     Phylogeny of mycoplasmalike organisms (phytoplasmas):
a basis for
            their classification
  JOURNAL   J. Bacteriol. 176 (17), 5244-5254 (1994)
   PUBMED   8071198
COMMENT     Original source text: Paulownia witches'-broom
mycoplasma-like
            organism DNA.
FEATURES             Location/Qualifiers
     source          1. .339
                     /organism="Paulownia witches'-broom
phytoplasma"
                     /mol_type="genomic DNA"
                     /db_xref="taxon:39647"
                     /lab_host="Catharanthus roseus"
ORIGIN

  Query Match             100.0%;  Score 21;  DB 15;  Length 339;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches   21;  Conservative    0;  Mismatches    0;  Indels
0;  Gaps    0;

Qy          1 AATAAGCCCCGGCTAACTATG 21
              |||||||||||||||||||||
Db        314 AATAAGCCCCGGCTAACTATG 334


RESULT 5
MPLRGDLY01
LOCUS       MPLRGDLY01               341 bp    DNA     linear
BCT 21-APR-1995
DEFINITION  Lethal yellows of palms mycoplasmalike organism 16S
ribosomal RNA
            (16S rRNA) gene, partial sequence.
ACCESSION   L27030
VERSION     L27030.1  GI:780098
KEYWORDS    16S ribosomal RNA.
SEGMENT     1 of 2
SOURCE      Palm lethal yellowing phytoplasma
  ORGANISM  Palm lethal yellowing phytoplasma
            Bacteria; Firmicutes; Mollicutes; Acholeplasmatales;
            Acholeplasmataceae; Candidatus Phytoplasma.
REFERENCE   1  (bases 1 to 341)
  AUTHORS   Gundersen,D.E., Lee,I.M., Rehner,S.A., Davis,R.E. and
            Kingsbury,D.T.
  TITLE     Phylogeny of mycoplasmalike organisms (phytoplasmas):
```

a basis for
            their classification
  JOURNAL   J. Bacteriol. 176 (17), 5244-5254 (1994)
   PUBMED   8071198
COMMENT   Original source text: Palm lethal yellowing
mycoplasma-like
            organism DNA.
FEATURES   Location/Qualifiers
     source   1. .341
                     /organism="Palm lethal yellowing phytoplasma"
                     /mol_type="genomic DNA"
                     /db_xref="taxon:39646"
                     /lab_host="Cocos nucifera"
ORIGIN

  Query Match   100.0%; Score 21; DB 15; Length 341;
  Best Local Similarity   100.0%; Pred. No. 0;
  Matches   21; Conservative   0; Mismatches   0; Indels
0; Gaps   0;

Qy   1 AATAAGCCCCGGCTAACTATG 21
              |||||||||||||||||||||
Db   316 AATAAGCCCCGGCTAACTATG 336

RESULT 6
MPLRGDPW01
LOCUS   MPLRGDPW01   341 bp   DNA   linear
BCT 07-DEC-1994
DEFINITION Potato witches'-broom mycoplasmalike organism 16S
ribosomal RNA
            (16S rRNA) gene, partial sequence.
ACCESSION   L27039
VERSION   L27039.1 GI:598309
KEYWORDS   16S ribosomal RNA.
SEGMENT   1 of 2
SOURCE   Potato witches'-broom phytoplasma
  ORGANISM Potato witches'-broom phytoplasma
            Bacteria; Firmicutes; Mollicutes; Acholeplasmatales;
            Acholeplasmataceae; Candidatus Phytoplasma; 16SrVI
(Clover
            proliferation group).
REFERENCE   1 (bases 1 to 341)
  AUTHORS   Gundersen,D.E., Lee,I.M., Rehner,S.A., Davis,R.E. and
            Kingsbury,D.T.
  TITLE   Phylogeny of mycoplasmalike organisms (phytoplasmas):

a basis for
their classification
JOURNAL J. Bacteriol. 176 (17), 5244-5254 (1994)
PUBMED 8071198
COMMENT Original source text: Potato witches'-broom
mycoplasma-like
organism DNA.
FEATURES Location/Qualifiers
source 1. .341
/organism="Potato witches'-broom phytoplasma"
/mol_type="genomic DNA"
/db_xref="taxon:37701"
/lab_host="Catharanthus roseus"
ORIGIN

Query Match 100.0%; Score 21; DB 15; Length 341;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 21; Conservative 0; Mismatches 0; Indels
0; Gaps 0;

Qy 1 AATAAGCCCCGGCTAACTATG 21
||||||||||||||||||||
Db 316 AATAAGCCCCGGCTAACTATG 336

RESULT 7
MPLRGDVR01
LOCUS MPLRGDVR01 341 bp DNA linear
BCT 07-DEC-1994
DEFINITION Beet leafhopper transmitted virescence mycoplasmalike
organism 16S
ribosomal RNA (16S rRNA) gene, partial sequence.
ACCESSION L27045
VERSION L27045.1 GI:598312
KEYWORDS 16S ribosomal RNA.
SEGMENT 1 of 2
SOURCE Beet leafhopper transmitted virescence phytoplasma
ORGANISM Beet leafhopper transmitted virescence phytoplasma
Bacteria; Firmicutes; Mollicutes; Acholeplasmatales;
Acholeplasmataceae; Candidatus Phytoplasma.
REFERENCE 1 (bases 1 to 341)
AUTHORS Gundersen,D.E., Lee,I.M., Rehner,S.A., Davis,R.E. and
Kingsbury,D.T.
TITLE Phylogeny of mycoplasmalike organisms (phytoplasmas):
a basis for
their classification

JOURNAL J. Bacteriol. 176 (17), 5244-5254 (1994)
PUBMED 8071198
COMMENT Original source text: Beet leafhopper transmitted virescence
mycoplasma-like organism DNA.
FEATURES Location/Qualifiers
source 1. .341
/organism="Beet leafhopper transmitted virescence
phytoplasma"
/mol_type="genomic DNA"
/db_xref="taxon:37694"
/lab_host="Catharanthus roseus"
ORIGIN

Query Match 100.0%; Score 21; DB 15; Length 341;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 21; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AATAAGCCCCGGCTAACTATG 21
||||||||||||||||||||
Db 316 AATAAGCCCCGGCTAACTATG 336

RESULT 8
AY389819
LOCUS AY389819 424 bp DNA linear BCT 18-JAN-2004
DEFINITION Aster yellows phytoplasma strain AY-SG 16S ribosomal RNA gene,
partial sequence.
ACCESSION AY389819
VERSION AY389819.1 GI:40807774
KEYWORDS .
SOURCE Aster yellows phytoplasma
ORGANISM Aster yellows phytoplasma
Bacteria; Firmicutes; Mollicutes; Acholeplasmatales; Acholeplasmataceae; Candidatus Phytoplasma; Candidatus Phytoplasma
asteris.
REFERENCE 1 (bases 1 to 424)
AUTHORS Zhang,J., Nault,L.R., Hogenhout,S.A., Hoy,C.W. and Miller,S.A.
TITLE Molecular and symptom analyses of phytoplasma strains collected